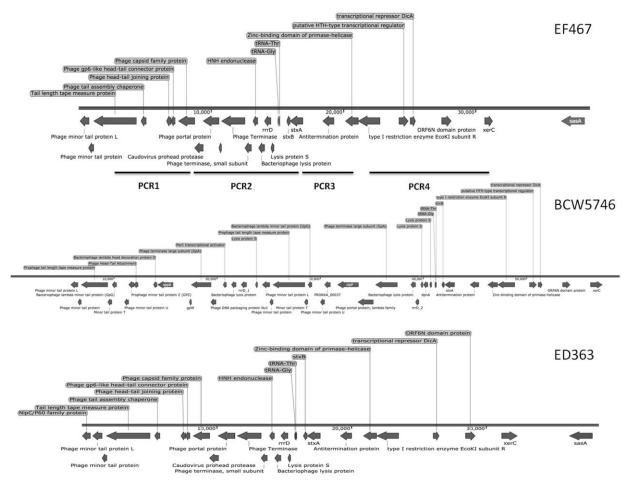
Whole Genome Characterization and Strain Comparison of VT2f-Producing *Escherichia* coli Causing Hemolytic Uremic Syndrome

Technical Appendix



Technical Appendix Figure. Maps of the partial VT2f phages constructed by using the whole-genome sequences of the verotoxigenic *Escherichia coli* strains EF467, isolated from fecal specimens from a patient with hemolytic uremic syndrome; BCW5746, isolated from fecal specimens from a patient with diarrhea; and ED363, isolated from feces specimens from pigeons. The long PCRs used to verify the phages' structures are indicated as black bars. orf, open reading frame.